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1648

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RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/581,976

DATE: 01/08/2001  
 TIME: 18:14:25

Input Set : A:\seqlist.txt  
 Output Set: N:\CRF3\01082001\I581976.raw

ENTERED

4 <110> APPLICANT: Dalemans, Wilfried L.J.  
 5 Gerard, Catherine Marie Ghislaine  
 7 <120> TITLE OF INVENTION: Vaccine  
 10 <130> FILE REFERENCE: B45124  
 12 <140> CURRENT APPLICATION NUMBER: 09/581,976  
 13 <141> CURRENT FILING DATE: 2000-06-20  
 15 <150> PRIOR APPLICATION NUMBER: PCT/EP98/08563  
 16 <151> PRIOR FILING DATE: 1998-12-18  
 18 <150> PRIOR APPLICATION NUMBER: GB 9727262.9  
 19 <151> PRIOR FILING DATE: 1997-12-24  
 21 <160> NUMBER OF SEQ ID NOS: 26  
 23 <170> SOFTWARE: FastSEQ for Windows Version 3.0  
 25 <210> SEQ ID NO: 1  
 26 <211> LENGTH: 220  
 27 <212> TYPE: PRT  
 28 <213> ORGANISM: Artificial Sequence  
 30 <220> FEATURE:  
 31 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus  
 32 influenza B and E7 from Human papilloma virus type  
 33 16)  
 35 <400> SEQUENCE: 1  
 36 Met Asp Pro Ser Ser His Ser Ser Asn Met Ala Asn Thr Gln Met Lys  
 37 1 5 10 15  
 38 Ser Asp Lys Ile Ile Ile Ala His Arg Gly Ala Ser Gly Tyr Leu Pro  
 39 20 25 30  
 40 Glu His Thr Leu Glu Ser Lys Ala Leu Ala Phe Ala Gln Ala Asp  
 41 35 40 45  
 42 Tyr Leu Glu Gln Asp Leu Ala Met Thr Lys Asp Gly Arg Leu Val Val  
 43 50 55 60  
 44 Ile His Asp His Phe Leu Asp Gly Leu Thr Asp Val Ala Lys Lys Phe  
 45 65 70 75 80  
 46 Pro His Arg His Arg Lys Asp Gly Arg Tyr Tyr Val Ile Asp Phe Thr  
 47 85 90 95  
 48 Leu Lys Glu Ile Gln Ser Leu Glu Met Thr Glu Asn Phe Glu Thr Met  
 49 100 105 110  
 50 Ala Met His Gly Asp Thr Pro Thr Leu His Glu Tyr Met Leu Asp Leu  
 51 115 120 125  
 52 Gln Pro Glu Thr Thr Asp Leu Tyr Cys Tyr Glu Gln Leu Asn Asp Ser  
 53 130 135 140  
 54 Ser Glu Glu Glu Asp Glu Ile Asp Gly Pro Ala Gly Gln Ala Glu Pro  
 55 145 150 155 160  
 56 Asp Arg Ala His Tyr Asn Ile Val Thr Phe Cys Cys Lys Cys Asp Ser  
 57 165 170 175  
 58 Thr Leu Arg Leu Cys Val Gln Ser Thr His Val Asp Ile Arg Thr Leu  
 59 180 185 190  
 60 Glu Asp Leu Leu Met Gly Thr Leu Gly Ile Val Cys Pro Ile Cys Ser  
 61 195 200 205

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62 Gln Lys Pro Thr Ser Gly His His His His His His
63      210      215      220
65 <210> SEQ ID NO: 2
66 <211> LENGTH: 663
67 <212> TYPE: DNA
68 <213> ORGANISM: Artificial Sequence
70 <220> FEATURE:
71 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
72      influenza B and E7 from Human papilloma virus type
73      16)
75 <400> SEQUENCE: 2
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77 attattgctc accgtggtgc tagcgggttat ttaccagagc atacgttaga atctaaagca      120
78 cttgcggttg cacaacaggc tgattattta gagcaagatt tagcaatgac taaggatggt      180
79 cgttttagtg ttattcacga tcacttttta gatggcttga ctgatgttgc gaaaaaatc      240
80 ccacatcgtc atcgtaaaqa tggccggttac tatgtcatcg actttacott aaaagaaatt      300
81 caaagtttag aaatgacaga aaactttgaa accatggcca tgcattggaga tacacctaca      360
82 ttgatgaat atattgttaga ttggcaacca gagacaactg atctctactg ttatgagcaa      420
83 ttaaatgaca gctcagagga ggaggatgaa atagatggtc cagctggaca agcagaaccg      480
84 gacagagccc attacaatat tgaacctttt tgttgcaagt gtgactctac gcttcggttg      540
85 tgcgtacaaa gcacacacgt agacattcgt actttggaag acctgttaat gggcacacta      600
86 ggaattgtgt gcccatctg ttctcagaaa ccaactagtg gccaccatca ccattcccat      660
87 taa
89 <210> SEQ ID NO: 3
90 <211> LENGTH: 822
91 <212> TYPE: DNA
92 <213> ORGANISM: Artificial Sequence
94 <220> FEATURE:
95 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
96      influenza B and E6 from Human papilloma virus type
97      16)
99 <400> SEQUENCE: 3
C--> 100 atggatccaa gcagccattc atcaaatatg gcgaataccc aaatgaaatc agacaaaatc      60
101 attattgctc accgtggtgc tagcgggttat ttaccagagc atacgttaga atctaaagca      120
102 cttgcggttg cacaacaggc tgattattta gagcaagatt tagcaatgac taaggatggt      180
103 cgttttagtg ttattcacga tcacttttta gatggcttga ctgatgttgc gaaaaaatc      240
104 ccacatcgtc atcgtaaaqa tggccggttac tatgtcatcg actttacott aaaagaaatt      300
105 caaagtttag aaatgacaga aaactttgaa accatggcca tgtttcagga cccacaggag      360
106 cgaccagaaa agttaccaca gttatgcaca gagctgcaaa caactataca tgatataata      420
107 ttagaatgtg tgtactgcaa gcaacagtta ctgcgaactg aggtatatga ctttgccttt      480
108 cgggatttat gcatagtata tagagatggg aatccatatg ctgtatgtga taaatgttta      540
109 aagtttttatt ctaaaattag tgagtataga cattattgtt atagtttgta tggacaaca      600
110 ttagaacagc aatacaacaa accgttgtgt gatttgttaa ttagggtgat taactgtcaa      660
111 aagccactgt gtcctgaaga aaagcaaaga catctggaca aaaagcaaag attccataat      720
112 ataaggggtc ggtggaccgg tcgatgtatg tcttgttgca gatcatcaag aacacgtaga      780
113 gaaaccagc tgactagtgg ccaccatcac catcaccatt aa
116 <210> SEQ ID NO: 4
117 <211> LENGTH: 273
118 <212> TYPE: PRT

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119 <213> ORGANISM: Artificial Sequence
121 <220> FEATURE:
122 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
123     influenza B and E6 from Human papilloma virus type
124     16)
126 <400> SEQUENCE: 4
127 Met Asp Pro Ser Ser His Ser Ser Asn Met Ala Asn Thr Gln Met Lys
128   1             5             10             15
129 Ser Asp Lys Ile Ile Ala His Arg Gly Ala Ser Gly Tyr Leu Pro
130             20             25             30
131 Glu His Thr Leu Glu Ser Lys Ala Leu Ala Phe Ala Gln Gln Ala Asp
132             35             40             45
133 Tyr Leu Glu Gln Asp Leu Ala Met Thr Lys Asp Gly Arg Leu Val Val
134             50             55             60
135 Ile His Asp His Phe Leu Asp Gly Leu Thr Asp Val Ala Lys Lys Phe
136             65             70             75             80
137 Pro His Arg His Arg Lys Asp Gly Arg Tyr Tyr Val Ile Asp Phe Thr
138             85             90             95
139 Leu Lys Glu Ile Gln Ser Leu Glu Met Thr Glu Asn Phe Glu Thr Met
140             100            105            110
141 Ala Met Phe Gln Asp Pro Gln Glu Arg Pro Arg Lys Leu Pro Gln Leu
142             115            120            125
143 Cys Thr Glu Leu Gln Thr Thr Ile His Asp Ile Ile Leu Glu Cys Val
144             130            135            140
145 Tyr Cys Lys Gln Gln Leu Leu Arg Arg Glu Val Tyr Asp Phe Ala Phe
146             145            150            155            160
147 Arg Asp Leu Cys Ile Val Tyr Arg Asp Gly Asn Pro Tyr Ala Val Cys
148             165            170            175
149 Asp Lys Cys Leu Lys Phe Tyr Ser Lys Ile Ser Glu Tyr Arg His Tyr
150             180            185            190
151 Cys Tyr Ser Leu Tyr Gly Thr Thr Leu Glu Gln Gln Tyr Asn Lys Pro
152             195            200            205
153 Leu Cys Asp Leu Leu Ile Arg Cys Ile Asn Cys Gln Lys Pro Leu Cys
154             210            215            220
155 Pro Glu Glu Lys Gln Arg His Leu Asp Lys Lys Gln Arg Phe His Asn
156             225            230            235            240
157 Ile Arg Gly Arg Trp Thr Gly Arg Cys Met Ser Cys Cys Arg Ser Ser
158             245            250            255
159 Arg Thr Arg Arg Glu Thr Gln Leu Thr Ser Gly His His His His His
160             260            265            270
161 His
164 <210> SEQ ID NO: 5
165 <211> LENGTH: 1116
166 <212> TYPE: DNA
167 <213> ORGANISM: Artificial Sequence
169 <220> FEATURE:
170 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
171     influenza B and E6E7 fusion from Human papilloma
172     virus type 16)

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174 <400> SEQUENCE: 5
175 atggatccaa gcagccattc atcaaatatg gcgaataccc aaatgaaatc agacaaaatc      60
176 attattgctc accgtggtgc tagcggttat ttaccagagc atacgttaga atctaaagca      120
177 cttgcgtttg cacaacaggc tgattattta gagcaagatt tagcaatgac taaggatggt      180
178 cgttttagtg ttattcaaga tcaactttta gatggcttga ctgatgttgc gaaaaaatc      240
179 ccacatcgtc atcgtaaaag tggccgttac tatgtcatcg actttacctt aaagaaaatt      300
180 caaagttagt aaatgacaga aaactttgaa accatggcca tgtttcagga cccacaggag      360
181 cgacccagaa agttaccaca gttatgcaca gagctgcaaa caactatata tgatataata      420
182 ttagaatgtg tytactgcaa gcaacagtta ctgcgacgtg aggtatatga ctttgccttt      480
183 cgggatttat gcatagtata tagagatggg aatccatatg ctgtatgtga taaatgttta      540
184 aagttttatt ctaaaattag tgagtataga cattattgtt atagtittga tggacaaca      600
185 ttagaacagc aatacaacaa accgttggtg gatctgttaa ttaggtgtat taactgtcaa      660
186 aagccaatgt gtccatgaag aaagcaaaag catctggaca aaaagcauag attccataat      720
187 ataaggggtc ggtggaccgg tcgatgtatg tcttggttga gatcatcaag aacacgtaga      780
188 gaaacccagc tgatgcattg agatacacct acattgcatt aatatatgtt agatttgcaa      840
189 ccagagacaa ctgatctcta ctgttatgag caattaaatg acagctcaga ggaggaggat      900
190 gaaatagatg gtccagctgg acaagcagaa ccggacagag cccattacaa tattgtaacc      960
191 ttttggttga agtggtactc tacgcttcgg ttgtgcgtac aaagcacaca cgtagacatt      1020
192 cgtactttgg aagacctgtt aatgggcaca ctaggaattg tytgcccatc ctgttctcag      1080
193 aaaccaacta gtggccaaca tcaccatcac cattaa      1116
195 <210> SEQ ID NO: 6
196 <211> LENGTH: 371
197 <212> TYPE: PRT
198 <213> ORGANISM: Artificial Sequence
200 <220> FEATURE:
201 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
202   influenza B and E6E7 fusion from Human papilloma
203   virus type 16)
205 <400> SEQUENCE: 6
206 Met Asp Pro Ser Ser His Ser Ser Asn Met Ala Asn Thr Gln Met Lys
207   1           5           10           15
208 Ser Asp Lys Ile Ile Ile Ala His Arg Gly Ala Ser Gly Tyr Leu Pro
209   20          25          30
210 Glu His Thr Leu Glu Ser Lys Ala Leu Ala Phe Ala Gln Gln Ala Asp
211   35          40          45
212 Tyr Leu Glu Gln Asp Leu Ala Met Thr Lys Asp Gly Arg Leu Val Val
213   50          55          60
214 Ile His Asp His Phe Leu Asp Gly Leu Thr Asp Val Ala Lys Lys Phe
215   65          70          75          80
216 Pro His Arg His Arg Lys Asp Gly Arg Tyr Tyr Val Ile Asp Phe Thr
217   85          90          95
218 Leu Lys Glu Ile Gln Ser Leu Glu Met Thr Glu Asn Phe Glu Thr Met
219   100         105         110
220 Ala Met Phe Gln Asp Pro Gln Glu Arg Pro Arg Lys Leu Pro Gln Leu
221   115         120         125
222 Cys Thr Glu Leu Gln Thr Thr Ile His Asp Ile Ile Leu Glu Cys Val
223   130         135         140
224 Tyr Cys Lys Gln Gln Leu Leu Arg Arg Glu Val Tyr Asp Phe Ala Phe
225   145         150         155         160

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DATE: 01/08/2001

TIME: 18:14:25

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Output Set: N:\CRF3\01082001\I581976.raw

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226 Arg Asp Leu Cys ile Val Tyr Arg Asp Gly Asn Pro Tyr Ala Val Cys
227                               165                               170                               175
228 Asp Lys Cys Leu Lys Phe Tyr Ser Lys ile Ser Glu Tyr Arg His Tyr
229                               180                               185                               190
230 Cys Tyr Ser Leu Tyr Gly Thr Thr Leu Glu Gln Gln Tyr Asn Lys Pro
231                               195                               200                               205
232 Leu Cys Asp Leu Leu ile Arg Cys ile Asn Cys Gln Lys Pro Leu Cys
233                               210                               215                               220
234 Pro Glu Glu Lys Gln Arg His Leu Asp Lys Lys Gln Arg Phe His Asn
235                               225                               230                               235                               240
236 ile Arg Gly Arg Trp Thr Gly Arg Cys Met Ser Cys Cys Arg Ser Ser
237                               245                               250                               255
238 Arg Thr Arg Arg Glu Thr Gln Leu Met His Gly Asp Thr Pro Thr Leu
239                               260                               265                               270
240 His Glu Tyr Met Leu Asp Leu Gln Pro Glu Thr Thr Asp Leu Tyr Cys
241                               275                               280                               285
242 Tyr Glu Gln Leu Asn Asp Ser Ser Glu Glu Glu Asp Glu ile Asp Gly
243                               290                               295                               300
244 Pro Ala Gly Gln Ala Glu Pro Asp Arg Ala His Tyr Asn ile Val Thr
245                               305                               310                               315                               320
246 Phe Cys Cys Lys Cys Asp Ser Thr Leu Arg Leu Cys Val Gln Ser Thr
247                               325                               330                               335
248 His Val Asp ile Arg Thr Leu Glu Asp Leu Leu Met Gly Thr Leu Gly
249                               340                               345                               350
250 ile Val Cys Pro ile Cys Ser Gln Lys Pro Thr Ser Gly His His His
251                               355                               360                               365
252 His His His
253                               370
255 <210> SEQ ID NO: 7
256 <211> LENGTH: 663
257 <212> TYPE: DNA
258 <213> ORGANISM: Artificial Sequence
260 <220> FEATURE:
261 <223> OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
262     influenza B and mutated E7 from Human papilloma
263     virus type 16)
265 <400> SEQUENCE: 7
266 atggatccaa gcagccattc atcaaatatg gcgaataccc aaatgaaatc agacaaaatc        60
267 attattgctc accgtgggtc tagcggttat ttaccagagc atacgttaga atctaaagca        120
268 cttgcgtttg cacaacaggc tgattattta gagcaagatt tagcaatgac taaggatggt        180
269 cgttttagtg ttattcacga tcacttttta gatggcttga ctgatgttgc gaaaaaatc        240
270 ccacatcgtc atcgtaaaga tggccgttac tatgtcatcg actttacctt aaaagaaatt        300
271 caaagtttag aaatgacaga aaactttgaa accatggcca tgcattgaga tacacctaca        360
272 ttgcatgaat atatgttaga ttgcaacca gagacaactg atctctacgg ttatcagcaa        420
273 ttaaatgaca gctcagagga ggaggatgaa atagatggtc cagctggaca agcagaaccg        480
274 gacagagccc attacaatat tgtaaccttt ttttgcaagt gtgactctac gcttcggttg        540
275 tgcgtacaaa gcacacacgt agacattcgt acttgggaag acctgttaut gggcacacta        600
276 ggaattgtgt gcccacatcg ttctcagaaa ccaactagtg gccaccatca ccacacccat        660
277 taa

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/581,976

DATE: 01/08/2001

TIME: 18:14:26

Input Set : A:\seqlist.txt

Output Set: N:\CRF3\01082001\I581976.raw

L:100 M:112 C: (48) String data converted to lower case,  
M:112 Repeated in SeqNo=3